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54 ATG	108 GGT G	162 TCA S	216 AAG K	270 ACC T	324 GCC A
GAC	GCA A	CCC	GAG	ACC	AAG K
CCA	GAG	TGG W	CCG P	${ m TTG}$	ACC
45 ACC	99 TGG W	153 CAC H	207 CCT P	261 CTC L	315 GGC G
ACG CGG M	CTG	AAA K	GAG	AAA K	CCA P
	CTG	GTC	GTG V	CCG	CTT L
36 GAG	90 GTG V	144 CAA Q	198 GTG V	252 AAG K	306 ATC I
ACA	GTT V	ATG M	CGT R	CAG Q	CCC
TCT	GTG V	AAG K	GCC	GTC	GGC
27 CTC	81 CTG L	135 ATC I	189 GGC G	243 CCT P	297 AGG R
GCA	AGC	CCT	TGG W	TTC	9 9
TGA	ACC	GTC V	GCC	CTG	CAG Q
18 CCC	72 GTC V	126 AAG K	180 AAG K	234 GTG V	288 GGT G
GCT	CTG	CCC	GAG E	GTG V	CGA R
GCA	CTC	GCA	CCA P	CTG	CCA P
9 ACT	63 CTC L	117 CCA P	171 GAC D	225 CAG Q	279 AAG K
200	AGG R	GTC V	CAG Q	GAC	GAG E
NNA	AGG R	GCA A	GAG	GAC D	GAG

IGURE 14

ACCUPANCE DESIGN

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378 CAT H	432 CGG R	486 GAC D	540 GTT	594 ATA
GAC	CCC	CAA Q	CCT	CAA
CCC	AGG R	GAC D	CGG	AGA
369 GAG E	423 GAG E	477 GAA E	531 CCC	585 GTT
CCC	GAG	GAG	TGN	CCA
AGT	9	CCG	NAG	TGC
360 CTG L	414 CAG Q	468 GGA G	22 AT	576 ACC
GTC V	GAC	CTG	2 3 3 6 6 7	CTA
CGT R	GAG	CTC	AGG	TTC
351 GGC G	405 GAG E	459 GTG V	513 TTC	567 ACC
CTG L	NCT	CAG Q	GGN 1	TGG
ACC	CCG	CAC H	TAG	GAT
342 GAC D	396 CCT P	450 AAT N	504 CAG Q	558 TING
GAG E	CAC H	CCA P	CCC	TGT
ACC	TAC	ATG M	CAA Q	AGG
333 GAG E	387 CTG L	441 GTG V	495 TAC Y	549 CCC
ATG M	AGC	TGG W	ATIN	AGG
TGG W	GAC D	TTG	CAC H	CCA

FIGURE 1B

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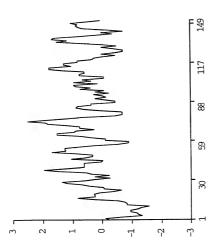
\vdash	MRRLLLLVTSLVVVLLWEAGAVPAPKVPIKM 2235738 MRRFLLATCLVAALLWEAGARPAHQVKT GI 899433
31	<u> QVKHWPSEQDPEKAWGARVVEPPEKDDQLV </u> 2235738 KG KHVFPEQETEKVWDTRALEPLEKDNQLG GI 899433
61	VLFPVQKPKLLTTEEKPRGQGRGPILPGTK 2235738 PLLLPEPKQKPAAAEEKRPDAM GI 899433
28	AWMETEDILGRVLSPEPDHDSLYHPP 2235738 TWVETEDILSHLRSPLQGPELDLDSIDHPM GI 899433
117	X E E D Q G B E R P R L W V M P N H Q V L L G P E E D Q D H 2235738 S D D V Q D E E V P Q S R P I L Y R Q V L Q G P E E D L D H GI 899433
147 142	$\begin{array}{c c} x \underline{Y Q P Q} \\ \underline{L A H S M E D S} \end{array} \hspace{1cm} 2235738$

FIGURE:

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IGURE 31